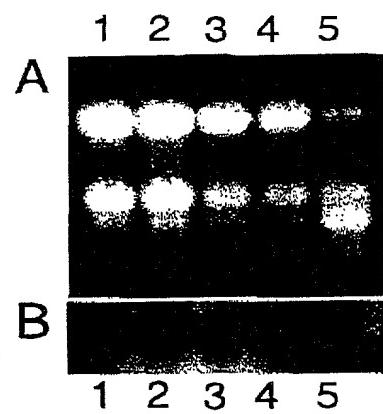


FIGURE 1

CGAAAAGAAAAATGGCTTGAATTAAATGCCATGCCCTGAAATCTCAGAAGCTCCCT 60  
M A L N F N A I A S K S Q K L P  
TGCTTTGCTCTTCCACCAAAGGCCACCCCTAGATCTCCCAAGTTCCATGATCTCCACC 120  
C F A L P P K A T L R S P K F S M I S T  
ATTCCTTCTGGCTCCAAAGAGGTGGAACTGAAAAAGCCTTCACGCCTCCAAAGGAG 180  
I P S G S K E V G N L K K P F T P P K E  
GTGCCTGTTAGATCACCCACTCCATGCCGCTCACAGAGATTGAGATCTTAAATCTTG 240  
V P V Q I T H S M P P H K I E I F K S L  
GAGGGCTGGCTGAGAACACATCTGACTCACCTCAAACCAGTTGAGAAATGTTGGCAA 300  
E G W A E N N I L T H L K P V E K C W Q  
CCCGCCGACTTCTTCAGATCTTAATTCTGATGGATTTCATGAGCAAGTCAAAGAGCTT 360  
P A D F L P D P N S D G F H E Q V K E L  
AGGGAAAGGGCAAAGGGAGATCCCAGATGATTACTTGTAGTTGGTGGTATATGATC 420  
R E R A K E I P D D Y F V V L V G D M I  
ACCGAGGAAGCCCTTCAACTTATCAAACAATGCTTAATACCTGGATGGAACCTCGTGAT 480  
T E E A L S T Y Q T M L N T L D G T R D  
GAGACAGGTGCTAGCCTTACCCCTGGGCCATTGGACCAGGGCTGGACTGCTGAAGAA 540  
E T G A S L T P W A I W T R A W T A E E  
AACAGGCATGGTGTGATCTGCTTAATAAGTATCTACTTGTCTGGGAGAGTGGACATGAGG 600  
N R H G D L L N K Y L Y L S G R V D M R  
CAAATTGAGAGGACAATCCAGTACTGATTGGATCGGAATGGATCCTCATACAGAGAAAT 660  
Q I E R T I Q Y L I G S G M D P H T E N  
AGTCCTTACCGAGGATTCAATATACTTCGTTCCAAGAAAGGGCAACTTTATTCCT 720  
S P Y R G F I Y T S F Q E R A T F I S H  
GGGAATACAGGCAGGCTGGCTAAGGAGTATGGGATATTAACCTGGCTAAATTGTGGT 780  
G N T G R L A K E Y G D I N L A Q I C G  
AGCATTGCCCTCAGATGAGAACGCCACGAGACAGCCTACACAAATCGTTGAAAAGCTG 840  
S I A S D E K R H E T A Y T K I V E K L  
TTTGAGATTGATCCTGATGAAACAGTCCTGGCATTGCTGACATGATGAAGAAGAAAATC 900  
F E I D P D E T V L A F A D M M K K I  
GCCATGCCGGCTGAGTCATCTATGATGGCAGAGATTAACTTATTTGACCACTACTCA 960  
A M P A E F I Y D G R D Y N L F D H Y S  
GCTGTTGCCAAAGAACATGGGTTTACACTGCTAAGGACTATGTTGATATAGTAGAGCAC 1020  
A V A Q R I G V Y T A K D Y V D I V E H  
CTGGTGGATCGATGGAAGGTGAAGGGAGCTAGCTGGCTTCAGGCCAGGGCGTAAAGCT 1080  
L V D R W K V K E L A G L S A E G R K A  
CAGGACTACTTGTGTTCACTTCCCTCGAGAATTAGAAGGTTAGAGGAGAGAGCGCAAGAA 1140  
Q D Y L C S L P S R I T R R L E E R A Q E  
AAGGCCAAGGAAGCACCAGTGTCCCATTCAAGTTGGATATTTGATAGAGAACGAAACTT 1200  
K A K E A P S V P F S W I F D R E V K L  
TAGGTGATGAAATACAGTTAACAGTCAATGCAATTGAGGAAACAAACACGAAGAAC 1260  
\*  
CTGAATGCCAACTTCTTTATATATCCGATGTAATAGAGGTTGTATATGTAACAGGAGG 1320  
AATTGCGTGGCTTGGTTAGGGTAGCAGCACATGTTCTGGATGTGTTGTCCTTAAAAAA 1380  
TAATGCCGATAGCGGCAGCTGTGATAGTTAGATGTTGTTTCAATGTCTGTTATA 1440  
TCGTTGAGGAGTAGTATGTTGTTGAAACAAATCTCATATCTTAGTGTATAA 1500  
TGATAATGCTGTGATGCAAGTTAGTTGCAATAAAAAAAAAAAAAAA 1553

FIGURE 2



**FIGURE 3**

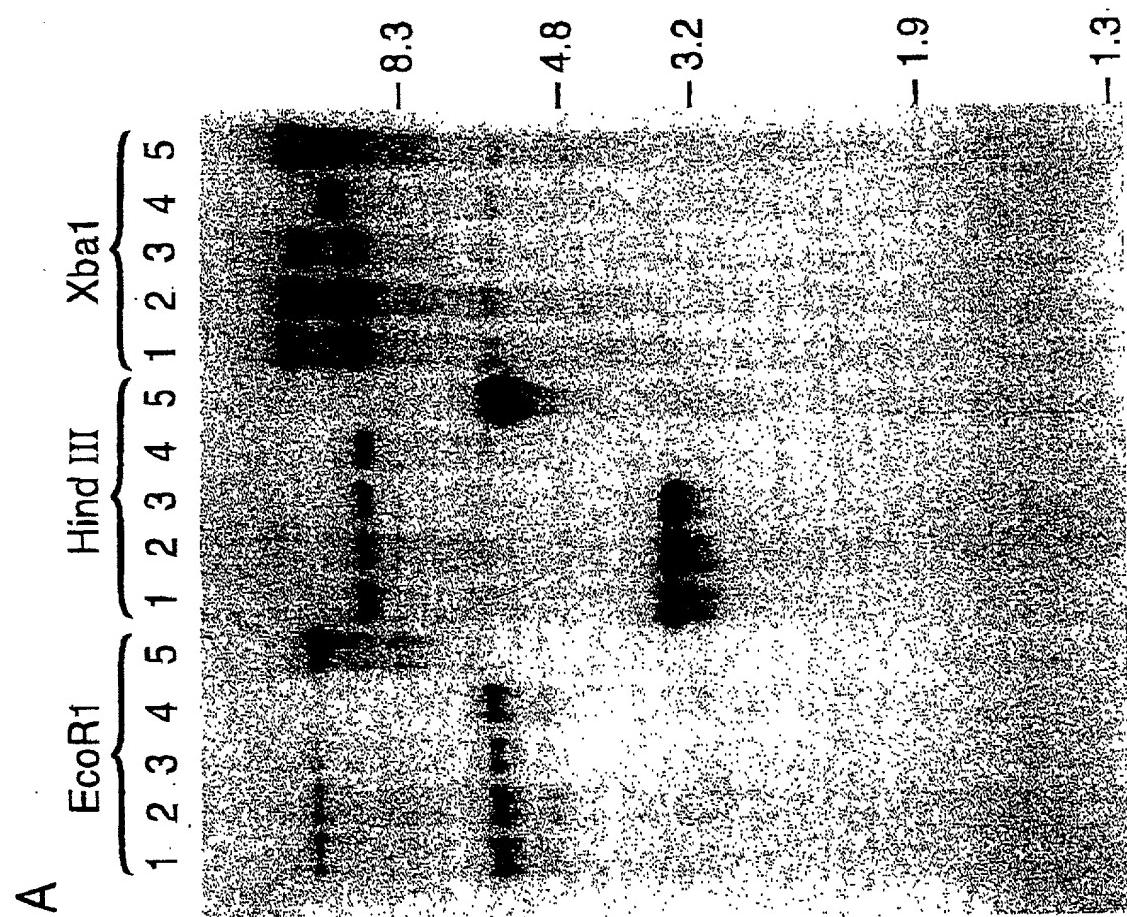


FIGURE 4A

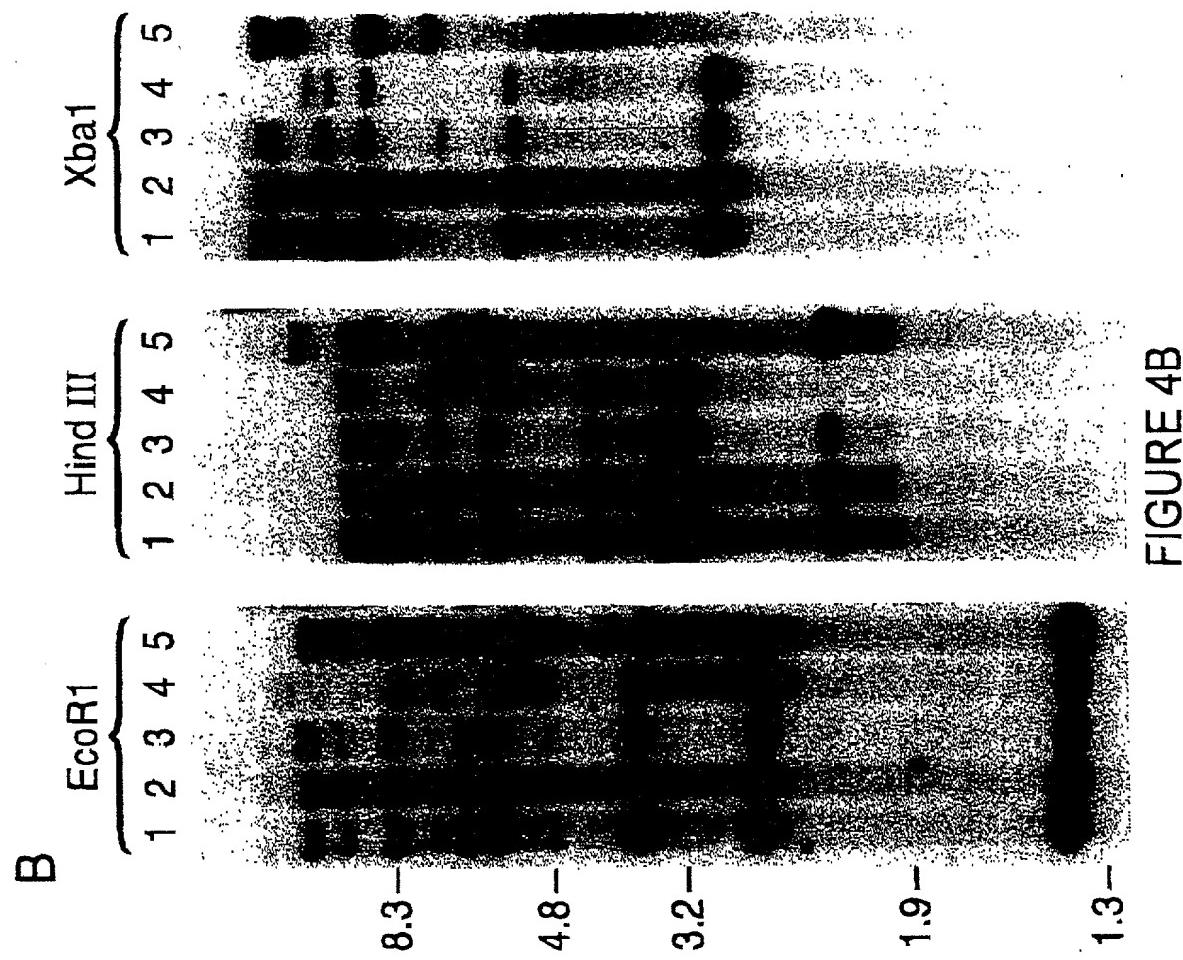


FIGURE 4B

CTCGCCCAAAACCAACACGCCCTCTTGCCCTCGTGTTCATCACCTGGCGTTAAACTGCT	60
TTCTTTAAAGCCAGAAAATGGGTGCCGGTGGTAGGATGCCATTGACGGTATAAAGGAG	120
M G A G G R M P I D G I K E	
GAAAATCGAGGCTCGGTCAATCGACTTCCGATCGAGAACGCCCGTTACGCTCGGTCAAG	180
E N R G S V N R V P I E K P P F T L G Q	
ATCAANGCAAGCCATTCCGCCCACTGTTTCCGCGCTCCCTCGATCCTCTCCTAC	240
I K Q A I P P H C F R R S L L R S F S Y	
<u>GTGGTCCATGACCTATGCTTAGCCTCTTACTACATTGCAACATCATATTTCAC</u>	300
<u>d12A4 primer</u>	
V V H D L C L A S F F Y Y I A T S Y F H	
TTTCTCCCACAACCCCTTTCTACATTGCTGGCCTGCTATTGGGTCTCCAAGGTTGC	360
F L P Q P F S Y I A W P V Y W V L Q G C	
ATCCTCACCGGTGTTGGGTCAATCGCACACGAGTGGGGTCAACCACGCTTCAGAGACTAC	420
I L T G V W V I A H E W G H H A F R D Y	
CAATGGGTTGACGACACCGTCGGGTTGATCCTCATTCCGCCCTTTAGTCCCGTACTTC	480
Q W V D D T V G L I L H S A L L V P Y F	
TCTGGAAAATCAGTCACCGCCGTCAACACTCGAACACCGGTTCCATGGAGCGTGACGAA	540
S W K I S H R R H S N T G S M E R D E	
GTATTCTGCCCAAACCCAAGTCTAAATTATCATGCTTGCAGAAATACCTAAACAATCCA	600
V F V P K P K S K L S C F A K Y L N N P	
CCCGGTGAGTTCTATCTCTGTAGTCACATTGACTCTGGTGGCTATGACTTAGCC	660
P G R V L S L V V T L T L G W P M Y L A	
TTCAACGTTGGGTGCGATACTATGATCGATTAGCTCCCACTATAACCCCTATGGCCCC	720
F N V S G R Y Y D R L A S H Y N P Y G P	
ATTTACTCCGATCGCGAGAGGCTACAAGTTACATCTCGATACTGGTATATTGCGGTA	780
I Y S D R E R L Q V Y I S D T G I F A V	
ATTTATGTACTTTATAAGATTGCTGCAACAAAAGGGCTGGCTTTATGCACTTAT	840
I Y V L Y K I A A T K G L A W L L C T Y	
GGGGTGCCTCTACTTATTGTGAATGCCCTCTGTGATCACCTACTTGCAACATACT	900
G V P L L I V N A F L V L I T Y L Q H T	
CACTCGGCATTGCCGCATTATGACTCGTCCGAATGGGATTGGTTGCGAGGAGCATTGCG	960
H S A L P H Y D S S E W D W L R G A L S	
ACGATGGATCGAGATTGGGGTGTGAACAAAGTGTCCATAACATCACCGATAACGCA	1020
T M D R D F G V L N K V F H N I T D T H	
GTTGCTCATCACCTCTCAACGATGCCACATTATCATGCAATGGAGGCCACTAAAGCA	1080
V A H H L F S T M P H Y H A M E A T K A	
ATCAAACCAACTCGGCAAGTATTACCTTCGACGGGACACCAGTTACAAGGCAATG	1140
I K P I L G K Y Y P F D G T P I Y K A M	
TGGAGGGAGGCAAAGAGTGCCTTACGTTGAGCCTGACGTTGGTGGTGGTGGTGGT	1200
W R E A K E C L Y V E P D V G G G G G	
AGCAAAGGTGTTTGGTATGTAACAAGTTCTAAAGACCGACCAACTGCCTGATAGCT	1260
S K G V F W Y R N K F *	
GGCCGGCGAAATCAACGTAACGTAAGTACTTATTAGACTAGTGTAACTAGGAAAGTTAATA	1320
ATTAATGGTAGGAAAATGTGGAATAGTGCCTAGTAGTTATGATTAAGTGTGTATT	1380
AATAAACTATATGGTAGAAAAAAAAAAAAAA	1411

FIGURE 5

taaaaaaaaaaaaggcattttcatcttaaagagacagcgaggaagccacgaagataata  
gagtgatttcaatctccattnaagggtgtggacaatgggtgctggaggcagaatgtc  
M G A G G R M S  
ggttccaaacgagtcacaaaaaccgaattcaactcaactgaagcgagttccatactcaaa  
V P T S P K K P E F N S L K R V P Y S K  
gcaccccttcactctgagtgaaatcaagaaagccatcccaccacactgttccagcgctc  
P P F T L S E I K K A I P P H C F Q R S  
cgttttacgctcatctcatatctcccttacgacttatattggcctcttttttacca  
V L R S F S Y L L Y D F I L A S L F Y H  
tgtggccaccaattactccctaacccttcaggctctccaacgtggcttggcctct  
V A T N Y F P N L P Q A L S N V A W P L  
ttattggccatgcaagggtgcattttgaccggcggttgggtcatagcccatgaatgtgg  
Y W A M Q G C I L T G V W V I A H E C G  
ccaccatgcttcagtgattatcaatggcttgacgacaccgtggccttacccactc  
H H A F S D Y Q W L D D T V G L I L H S  
ttctctcttagtccatattcttggaaatatagccaccggcgtcaccattctaacac  
S L L V P Y F S W K Y S H R R H H S N T  
cggtccctcgaaaggatgaagtgtcggtccaaagaaaaatctggttaaagatggtg  
G S L E R D E V F V P K K K S G L R W W  
ggccaaacacttcaacaatccacgggtcggttctgtcaatcaccattcaacttaccct  
A K H F N N P P G R F L S I T I Q L T L  
tggttggccgcttacttagcttcaacgttgccggccggccttacgacaggtcgctg  
G W P L Y L A F N V A G R P Y D R F A C  
ccactatgacccttacggccccatatttccgaccgggaacgactccaaatctatatctc  
H Y D P Y G P I F S D R E R L Q I Y I S  
tgacgcccggcgtcccgctgtcgccctatgcgcctaccgtctcggttggccaaagggg  
D A G V L A V A Y A L Y R L V L A K G V  
aggttgggttattagcgttatgggtgccattattgggttaacgcctttagtaat  
G W V I S V Y G V P L L V V N A F L V M  
gatcacgtatttgcacacactcaccatcttgcgcactatgattcctcgagtg  
I T Y L Q H T H P S L P H Y D S S E W D  
ctggatgagaggagcttatactgtggacagagattatgggattttaaacaagggttt  
W M R G A L S T V D R D Y G I L N K V F  
ccataacataaccgacactcatgtggctcatattgtttcgacaatgcctcaactatca  
H N I T D T H V A H H L F S T M P H Y H  
tgccatggggccaccaaggcgataaagccatattggggaaatactatcagttcgatgg  
A M V A T K A I K P I L G E Y Y Q F D G  
gatgcctgrctataaggcgatatggaggaggcgaaaggagtgtctacgttgcaccaga  
M P V Y K A I W R E A K E C L Y V E P D  
tgaggcgacaaaggataaagggtgtgtttgggttagaaacaagctttaatattgcatt  
E G D K D K G V F W F R N K L \*  
ttaccttaggcatt  
taatggtagaaataaatacacacagcatggattggcaataaaaaa

FIGURE 6

gmfAD2 - 2	1	- - - - -	W GAG GRD DVEPPAN	HL LFGP
atfAD2	1	- - - - -	W GAG GRG RPPVETS	HL LQPF
ghfAD2 - 1	1	- - - - -	W GAG GRM PDDGIK	HL LQPF
gmfAD2 - 1	1	- - - - -	W OLAKET THG GRDAKE	HL LQPF
bnfAD1	1	- - - - -	W VVWVVAQDQ	HL LQPF
gmfAD6	1	MACTLADSLLEFFKGSYQKFVLLRDIARYSPGIFPSLNBNGLIQKARRQRNFVTRNKWTVTHAYATEFQV	HL LQPF	
gmfAD2 - 2	14	EVKSE EVD PPHN RVPPEKPEPCE	HL LQPF	
acfAD2	14	EVGDEKKAKIPPHCFIRSTPFSYEE	HL LQPF	
ghfAD2 - 1	14	EVGDEKKAKIPPHCFIRSTPFSYEE	HL LQPF	
gmfAD2 - 1	14	EENR. EENR. EENR. EENR.	HL LQPF	
bnfAD1	20	VQKR. KPESSRVPNTK PPF.	HL LQPF	
gmfAD6	9	EVNANDE FEDSAQEEF. KEGDIEAIPHKCWVLSPLSMSYVARD.	HL LQPF	
gmfAD6	71	APVEEYVOLADEYQFRQVGEPLSDDTLKDVXNPLPKREVYEDDKAWBSVLTISVTEYALGFLWISK	HL LQPF	

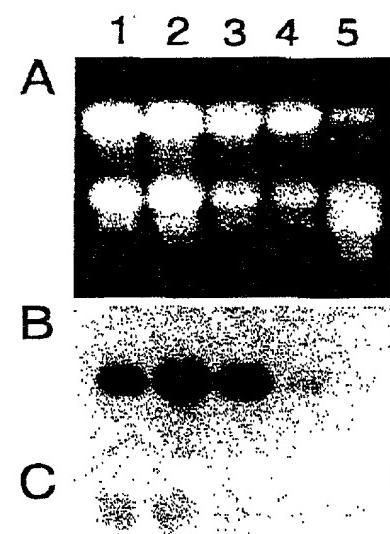
GmFAD2 - 2	61	L S A R O M A Y V A Q G C L T G V W V I A H E C H A F S D Y Q L D D T V G L L H S A L L V P V F S W K Y G H R R H H S N T G S
GtFAD2	61	L S A R P L Y V A Q G C L T G V W V I A H E C H A F S D Y Q L D D T V G L L H S P L L V P V F S W K Y G H R R H H S N T G S
gHfAD2 - 1	60	F S I A R P F V V W Q G C L T G V W V I A H E C H A F S D Y Q L D D T V G L L H S A L L V P V F S W K Y G H R R H H S N T G S
GmFAD2 - 1	65	T S I A R P E Y V V W Q G C L T G V W V I A H E C H A F S D Y Q L D D T V G L L H S T L L V P V F S W K Y G H R R H H S N T G S
bhFAD1	69	S I F F W P L Y V A Q D G T V V W Q G C L T G V W V I A H E C H A F S D Y Q L D D T V G L L H S T L L V P V F S W K Y G H R R H H S N T G S
gmyFAD6	141	A P M Y L L E P A N V W T G T A F G P V V I L H C C H A F S S N K L D D T V G L L H S T L L V P V F S W K Y G H R R H H S N T G S

GmFA D2 - 2 151 LERDEVFVKOKSKCKKQSKYLNWPEGRVTELAUTLGLWPPLYLAFNVSGRYDRAFTCHYDPYDPIVSDR  
ATFA D2 151 LERDEVFVKOKSKCKKQSKYLNWPEGRVTELVQFVLGWPPLYLAFNVSGRYDGCFAECHEFENP  
gHfA D2 - 1 150 LERDEVFVKPKSKCKKQSKYLNWPEGRVTELVQFVLGWPPLYLAFNVSGRYDRAASHYNPYCPYDPIVSDR  
gMfA D2 - 1 155 LDRADEVFVKPKSKCKKQSKYLNWPEGRVTELVQFVLGWPPLYLAFNVSGRYDFASSHYPYEPYDPIVSDR  
bnfA D1 138 MENDESAVPEPERLYKMLGKNSTRMAYTQIPMLOAEPLKREAPGNE...ESHYNPYEPYDPIVSDR  
gefA D6 211 LREDTAENE...VVKDEFEESTLLERKAING...VOPERCHWMJIAHWLNUWHEDLKKFPRPSIV

gmfad2 - 2 221 ERLQIVYNSDAGYLAVVYGLMRKANAKGLAVWVCCYGVPLLLVNQFLVLITLQH. THEALPHYTSSEDW  
 atfad2 221 ERLQIVYNSDAGYLAVVYGLMRKANAKGLAVWVCCYGVPLLLVNQFLVLITLQH.  
 ghpfd2 - 1 220 ERLQIVYNSDTGSEFAVVYVLLNATRGLAVLWCCYGVPLLLVNQFLVLITLQH.  
 gmfad2 - 1 225 ERLQIVYNSDVMFENFVYVLLNATRGLAVLWCCYGVPLLLVNQFLVLITLQH.  
 200 ERALIASSTQVSIQIATEVYVLLNATRGLAVLWCCYGVPLLLVNQFLVLITLQH.  
 bnfad3 267 PRYKEEACVFAE.  
 gmfad6

GMFA D2 - 2	289	WLR GAL . AT VDR DYG I LNK V F H N I T D T V A H H L F S T M P H Y H A M E A T K A I K P I L G E Y Y A F D S T P V K A M V R
ATFADA	289	WLR GAL . AT VDR DYG I LNK V F H N I T D T V A H H L F S T M P H Y H A M E A T K A I K P I L G Y P F D G T P V Y A M E R
GHDAD2 - 1	288	WLR GAL . AT VDR DYG I LNK V F H N I T D T V A H H L F S T M P H Y H A M E A T K A I K P I L G Y P F D G T P V Y A M E R
GMFD2 - 1	293	WNGAL . AT VDR DYG I LNK V F H N I T D T V A H H L F S T M P H Y H A M E A T K A I K P I L G Y P F D D T P V Y K A N W R
BNFAD3	270	WLRGAL . AT VDR DYG I LNK V F H N I T D T V A H H L F S T M P H Y H A M E A T K A I K P I L G Y P F D D T P V Y K A N W R
GMFA D6	334	RADGCL . AT VDR DYG I LNK V F H N I T D T V A H H L F S T M P H Y H A M E A T K A I K P I L G Y P F D D T P V Y K A N W R

gmpad2-2	158	EAECECLYVEPDDGEGE	...BKVQWVNNKL
ctfad2	159	EAECECLYVEPDEECK	...BKVQWVNNKL
gfad2-1	157	EAECECLYVEPDVGQGGGSGKVQWVNNKL	
gmfad2-1	162	EAECECLYVEPDGQGGGSGKVQWVNNKL	
bnpad2	136	VEELVAAEKKKDKYVSDDGDIVLYETDPPDLYVVAEDRSKIN	A
gmpad6	403	INTVGOVYKREYSLLCCLLRRTCP	



**FIGURE 8**

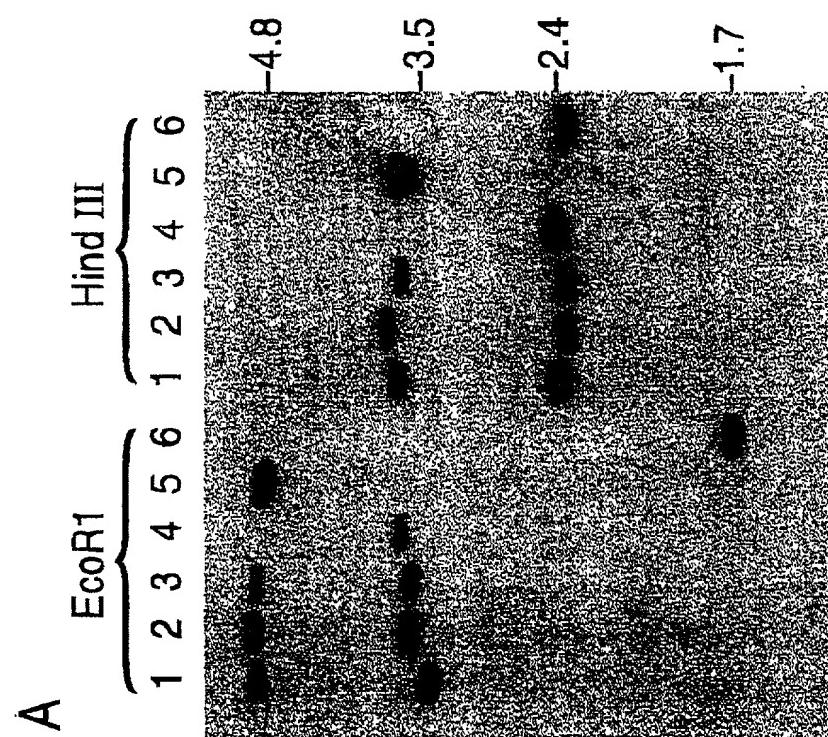


FIGURE 9A

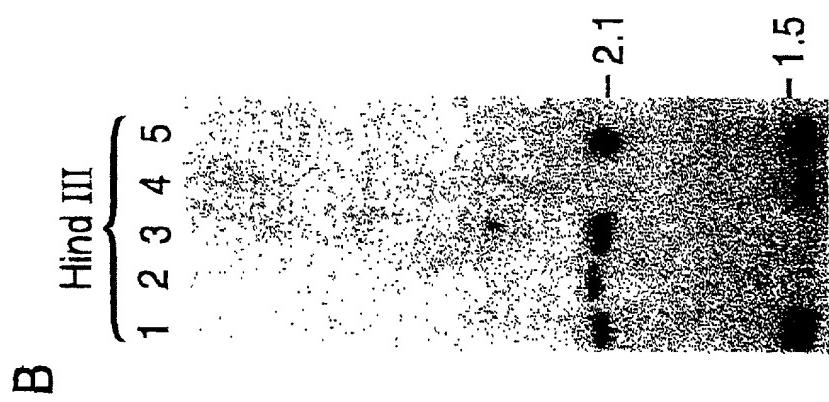


FIGURE 9B

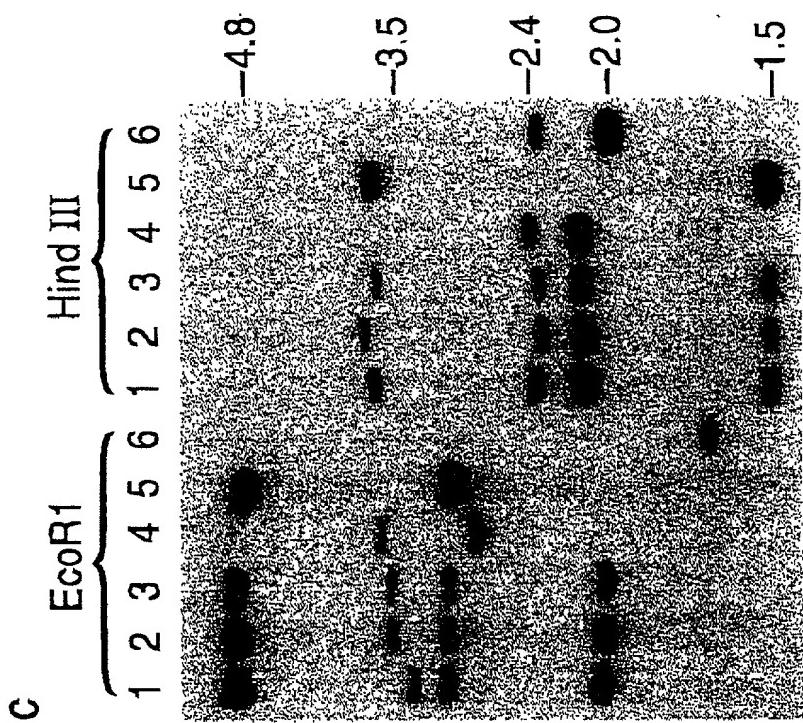


FIGURE 9C

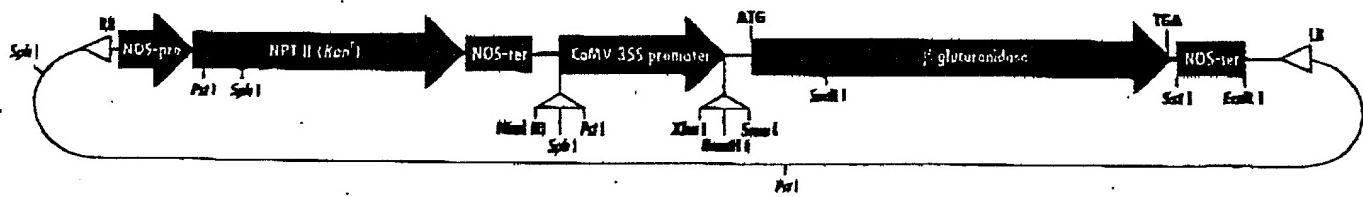
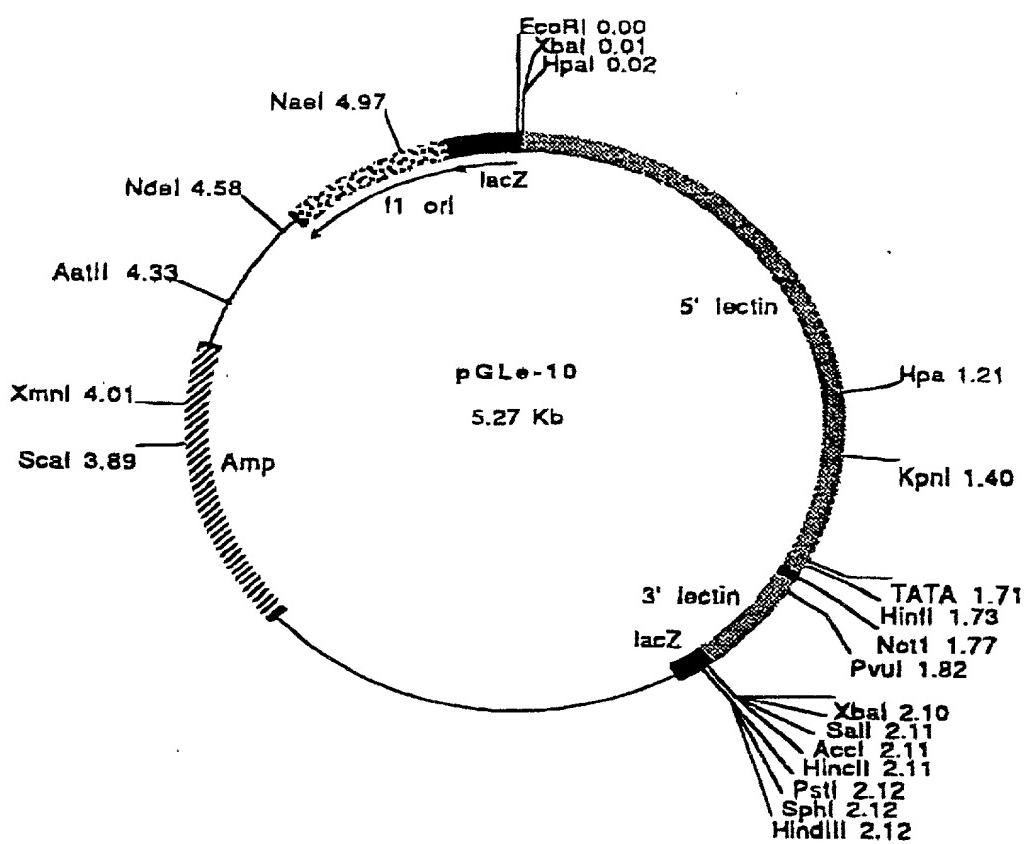


FIGURE 10

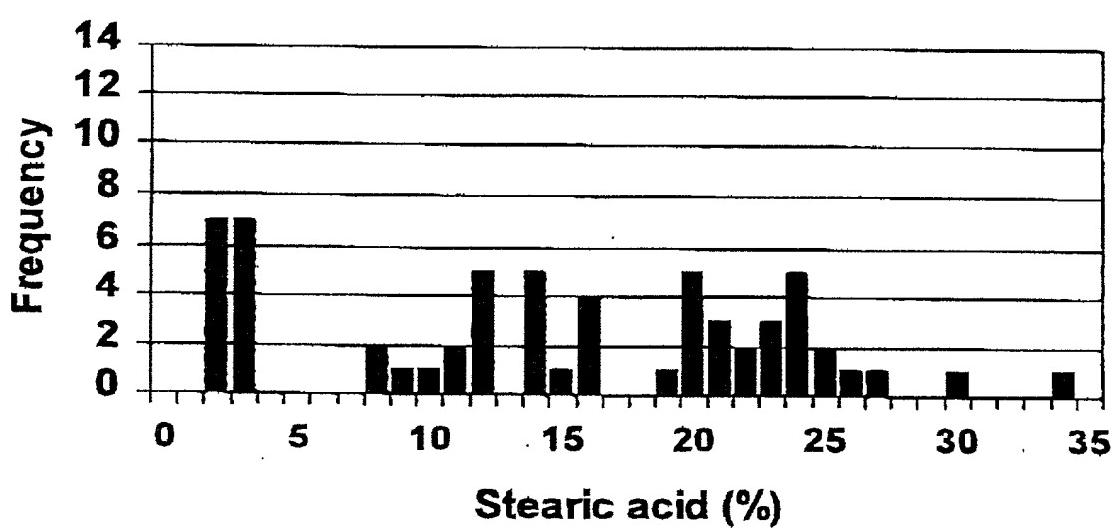
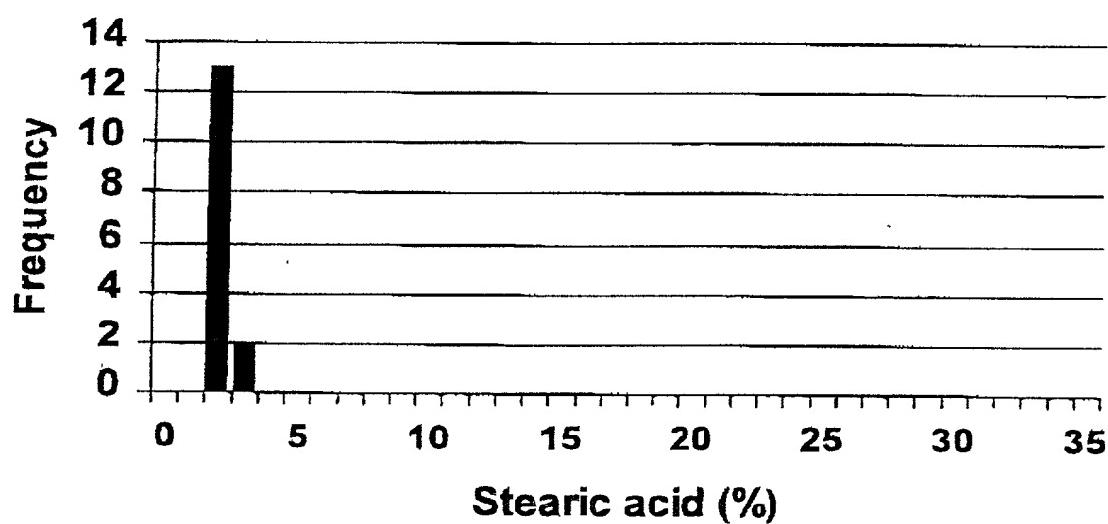


FIGURE 11

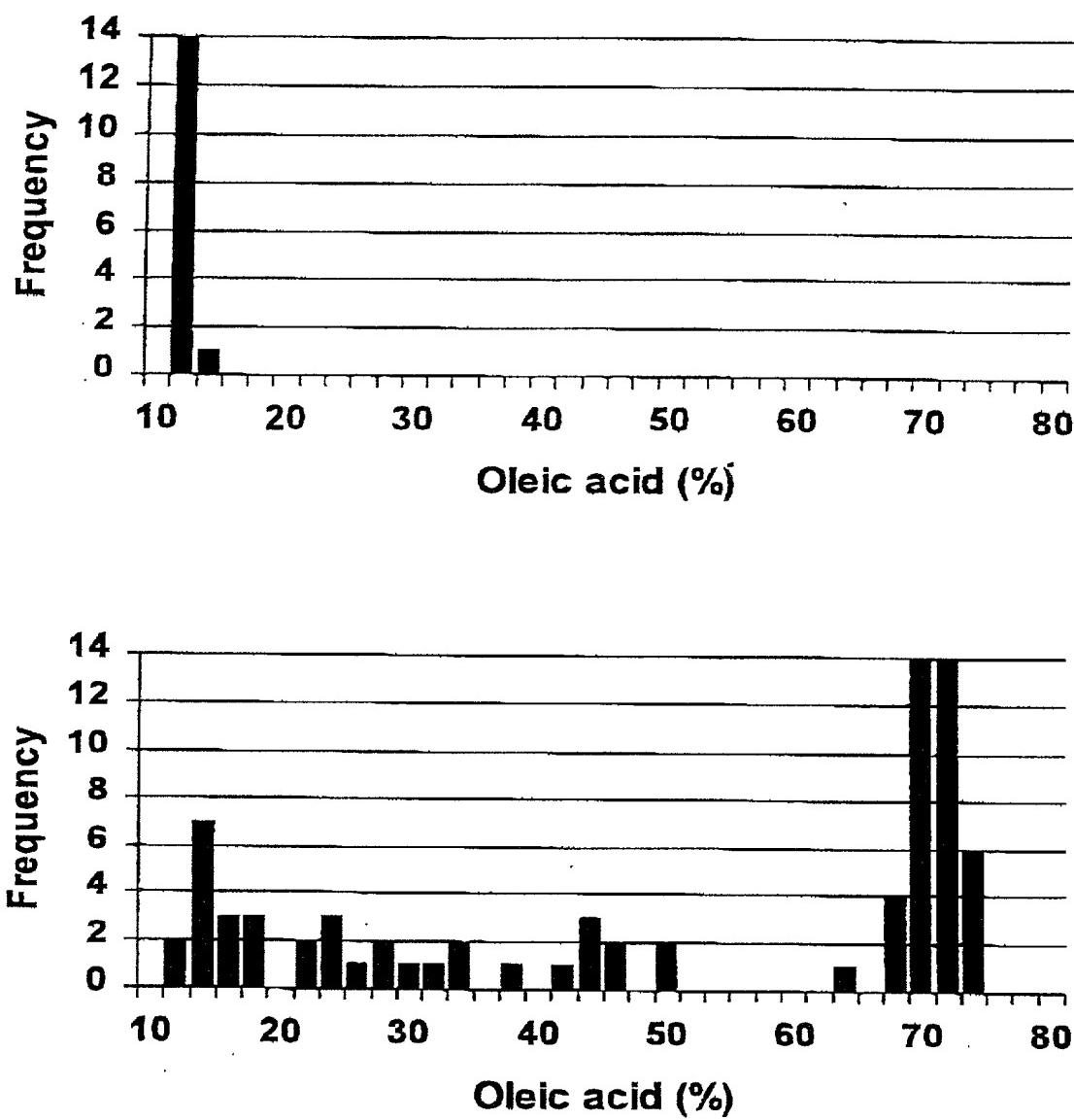


FIGURE 12

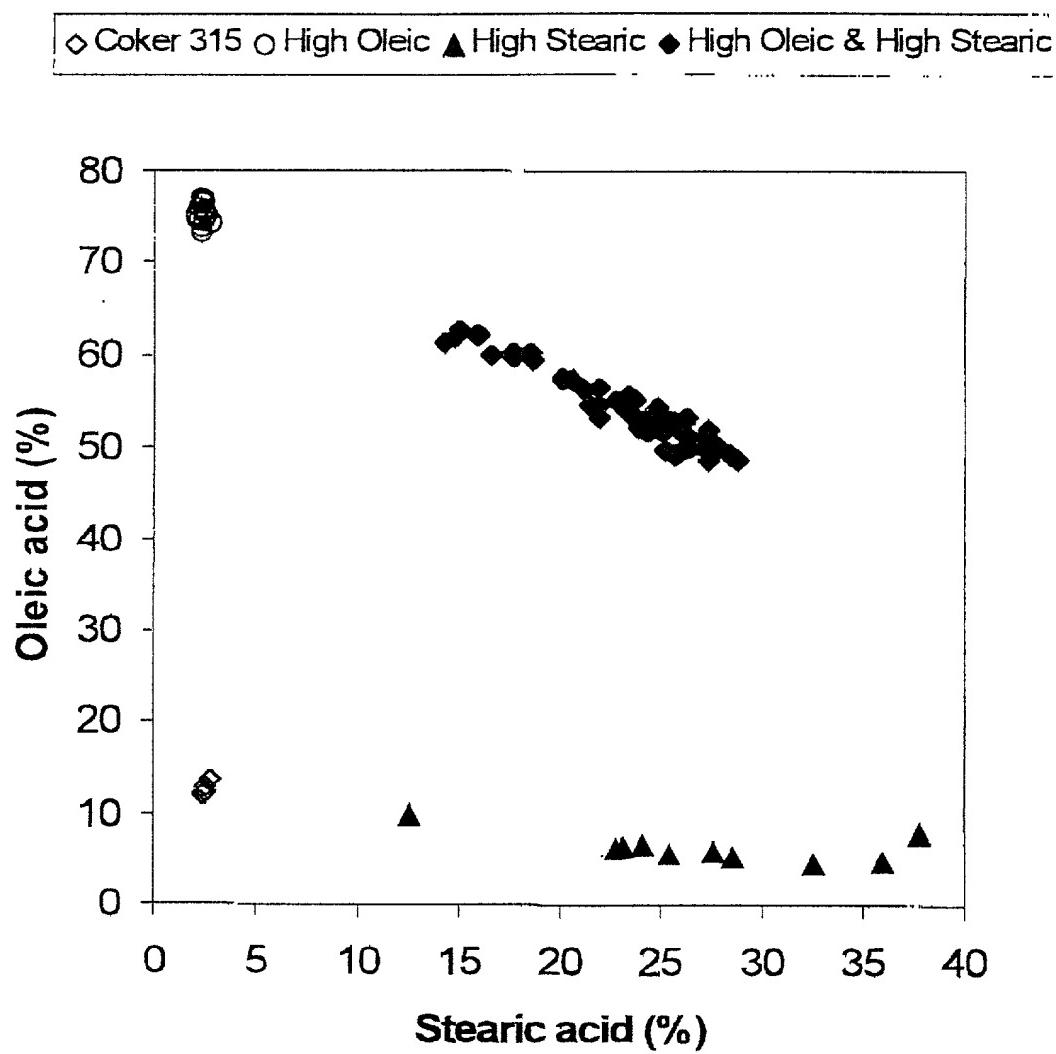


FIGURE 13